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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Minimum DB seq length: 0
Maximum DB seq length: 2000000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OM protein - protein search, using sw model
                 Published_Applications_AA:*

1: /cgn2_6/ptodata/2/pubpaa/US08_NEW_PUB.pep:*

2: /cgn2_6/ptodata/2/pubpaa/PCT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/2/pubpaa/US06_NEW_PUB.pep:*

4: /cgn2_6/ptodata/2/pubpaa/US06_PUBCOMB.pep:*

5: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

6: /cgn2_6/ptodata/2/pubpaa/US07_PUBCOMB.pep:*

7: /cgn2_6/ptodata/2/pubpaa/PCTUS.PUBCOMB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/US08_PUBCOMB.pep:*

9: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

10: /cgn2_6/ptodata/2/pubpaa/US09_NEW_PUB.pep:*

11: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

12: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

13: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*

13: /cgn2_6/ptodata/2/pubpaa/US10_NEW_PUB.pep:*
                      BLOSUM62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   188354 seqs, 42170167 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gapop 10.0 ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 MAEPGHSHHLSARVRGRTER.....LGRSNHLPPRGLLMDLTQCR 1001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
/cgn2_6/ptodata/2/pubpaa/US60_PUBCOMB.pep:*
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result	Score	Query Match	Length	DB	ID	Description
1	5341	97.0	1013	و	US-10-028-072-38	Sequence 38, Appl
2	5341	97.0	1013	ဖ	US-10-121-049-38	Sequence 38, Appl
ω	5341	97.0	1013	9	US-10-123-904-38	Sequence 38, Appl
4	5341	97.0	1013	9	US-10-140-470-38	Sequence 38, Appl
ഗ	5341	97.0	1013	9	US-10-175-746-38	•
σ	5341	97.0	1013	9	US-10-176-918-38	Sequence 38, Appl
7	5341	97.0	1013	9	US-10-176-921-38	Sequence 38, Appl
æ	5341	97.0	1013	9	US-10-137-865-38	Sequence 38, Appl
9	5341	97.0	1013	9	US-10-140-474-38	Sequence 38, Appl
10	5341	97.0	1013	9	US-10-142-431-38	Sequence 38, Appl
11	5341	97.0	1013	φ	US-10-143-114-38	Sequence 38, Appl
12	5341	97.0	1013	φ	US-10-140-002-38	Sequence 38, Appl
13	1448	26.3	464	9	US-10-002-050-20	Sequence 20, Appl
14	1448	26.3	464	9	US-10-002-304-20	Sequence 20, Appl
15	1448	26.3	464	13	US-10-003-152-20	e 20
16	1307.5	23.7	411	9	US-10-002-050-10	Sequence 10, Appl
17	1307.5	23.7	411	9	US-10-002-304-10	10,
18	1307.5	23.7	411	1	US-10-003-152-10	Sequence 10, Appl
19	889.5	16.2	208	9	US-09-925-299-982	Sequence 982, App

45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21	20
158	158	158	158	159	160.5	160.5	163	163	163	164	168	170	174	174	182	192	209.5	209.5	224	225	233	233	251	273	889.5
2.9	2.9	2.9	2.9	2.9	2.9	2.9	3.0	3.0	3.0	3.0	3.1	3.1	3.2	3.2	ω .ω	3.5	3.8	3. 8	4.1	4.1	4.2	4.2	4.6	5.0	16.2
1218	1218	1187	1036	1169	1193	1111	3084	2165	2150	610	50	1786	1786	1786	64	1587	3571	3571	60	1607	3594	3594	1609	81	208
9	9	10	10	10	10	10	10	10	10	9	10	10	10	10	10	10	10	9	10	10	10	9	10	10	10
US-10-219-247-7	US-10-219-248-7	US-09-995-593A-7	US-09-995-593A-6	US-09-801-368-106	US-09-756-071B-13	US-09-756-071B-15	US-09-938-275-4	US-09-800-729-155	US-09-321-987B-2	US-09-802-640-36	US-09-864-761-39644	US-09-938-275-7	US-09-938-275-6	US-09-873-676-113	US-09-864-761-47095	US-09-845-583-10	US-09-911-842-2	US-10-150-821-2	US-09-864-761-39057	US-09-938-275-10	US-09-911-842-4	US-10-150-821-4	US-09-938-275-11	US-09-864-761-39769	US-09-925-299-982
sequence /, Appli	, ;	Sequence /, Appli	Sequence 6, Appli	Sequence 106, App	Sequence 13, Appl	Sequence 15, Appl	Sequence 4, Appli	Sequence 155, App	Sequence 2, Appli	Sequence 36, Appl	Sequence 39644, A	Sequence 7, Appli	Sequence 6, Appli	Sequence 113, App	Sequence 47095, A	Sequence 10, Appl	Sequence 2, Appli	Sequence 2, Appli	Sequence 39057, A	Sequence 10, Appl	Sequence 4, Appli	Sequence 4, Appli	Sequence 11, Appl	Sequence 39769, A	

ALIGNMENTS

RESULT 1 US-10-028-072-38 US-10-028-072-38 Sequence 38, Application US/10028072 Publication No. US20030004311A1 FILE REFERENCE:

CURRENT APPLICATION NUMBER: US/10/028,072

CURRENT FILING DATE: 2001-12-19

PRIOR APPLICATION NUMBER: 60/049911

PRIOR FILING DATE: 1997-06-18

PRIOR APPLICATION NUMBER: 60/05974

PRIOR APPLICATION NUMBER: 60/059113

PRIOR APPLICATION NUMBER: 60/059113

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059117

PRIOR APPLICATION NUMBER: 60/059117

PRIOR FILING DATE: 1997-09-17

PRIOR FILING DATE: 1997-09-17

PRIOR APPLICATION NUMBER: 60/059122

PRIOR APPLICATION NUMBER: 60/059124

PRIOR APPLICATION NUMBER: 60/059124

PRIOR APPLICATION NUMBER: 60/059124

PRIOR APPLICATION NUMBER: 60/059127

PRIOR APPLICATION NUMBER: 60/059127

PRIOR APPLICATION NUMBER: 60/059184

PRIOR APPLICATION NUMBER: 60/059184 GENERAL INFORMATION: APPLICANT: APPLICANT:
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APPLICANT: APPLICANT:
APPLICANT: APPLICANT: APPLICANT: APPLICANT: Baker, Kevin P. APPLICANT: Beresini, Maur APPLICANT: Zhang TITLE OF INVENTION: APPLICANT: APPLICANT: Godowski, Paul J. Gurney, Austin L. Sherwood, Steven Watanabe, Colin K Wood, William Gerritsen, Mary E. Goddard, Audrey Desnoyers, Luc Filvaroff, Ellen DeForge, Laura Smith, Victoria Stewart, Timothy A. Gao, Wei-Qiang Beresini, Maureen Tumas,Daniel

PRIOR OR FILING DATE: 1998-01-23
DR APPLICATION NUMBER: 60/072320
DR FILING DATE: 1998-01-23
DR APPLICATION NUMBER: 60/073612
DR FILING DATE: 1998-02-04
DR APPLICATION NUMBER: 60/074086
DR FILING DATE: 1998-02-09
DR APPLICATION NUMBER: 60/074092
DR FILING DATE: 1998-02-09
DR APPLICATION NUMBER: 60/077791
DR APPLICATION NUMBER: 60/077791 OR APPLICATION NUMBER: 60/065186
OR FILING DATE: 1997-11-17
OR APPLICATION NUMBER: 60/065846
OR FILING DATE: 1997-11-17
OR APPLICATION NUMBER: 60/065846
OR FILING DATE: 1997-11-21
OR APPLICATION NUMBER: 60/066453
OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/066511
OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/066511
OR FILING DATE: 1997-11-24
OR APPLICATION NUMBER: 60/066770
OR FILING DATE: 1997-12-11
OR APPLICATION NUMBER: 60/069212
OR FILING DATE: 1997-12-11
OR APPLICATION NUMBER: 60/069218
OR APPLICATION NUMBER: 60/06934
OR APPLICATION NUMBER: 60/06934
OR APPLICATION NUMBER: 60/069694
OR APPLICATION NUMBER: 60/069694 OR FILING DATE: 1997-09-24
OR APPLICATION NUMBER: 60/062250
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/062285
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/06287
OR FILING DATE: 1997-10-17
OR APPLICATION NUMBER: 60/062814
OR FILING DATE: 1997-10-24
OR APPLICATION NUMBER: 60/062816
OR FILING DATE: 1997-10-24
OR FILING DATE: 1997-10-24 DR APPLICATION NUMBER: 60/063045
DR FILING DATE: 1997-10-24
DR APPLICATION NUMBER: 60/063082
DR FILING DATE: 1997-10-31
DR APPLICATION NUMBER: 60/063127
DR FILING DATE: 1997-10-24
DR FILING DATE: 1997-10-27
DR FILING DATE: 1997-10-27 OR FILING DATE: 1997-10-29

OR APPLICATION NUMBER: 60/063735

OR FILING DATE: 1997-10-29

OR APPLICATION NUMBER: 60/063738

OR FILING DATE: 1997-10-29

OR APPLICATION NUMBER: 60/063755

OR APPLICATION NUMBER: 60/063755 R APPLICATION NUMBER: 60, R FILING DATE: 1997-10-21 R PPLICATION NUMBER: 60, R FILING DATE: 1997-10-22 R APPLICATION NUMBER: 60, R APPLICATION NUMBE R APPLICATION NI R FILING DATE: R APPLICATION NI R FILING DATE: R APPLICATION NI APPLICATION NUMBER: 60/064248
FILING DATE: 1997-11-03
APPLICATION NUMBER: 60/064809
FILING DATE: 1997-11-07 APPLICATION NUMBER: 6 FILING DATE: 1997-10-NUMBER: 6 NUMBER: 60/059352 : 1997-09-19 NUMBER: 60/063733 60/063550 60/063704 60/063561 60/059836 60/059588 29

PRIOR OR APPLICATION NUMBER: 60/079728
OR FILING DATE: 1998-03-27
OR APPLICATION NUMBER: 60/080165
OR FILING DATE: 1998-03-31
OR APPLICATION NUMBER: 60/081203
OR FILING DATE: 1998-04-09
OR APPLICATION NUMBER: 60/081229
OR APPLICATION NUMBER: 60/081695
OR APPLICATION NUMBER: 60/081695
OR APPLICATION NUMBER: 60/081695 R APPLICATION NUMBER: R FILING DATE: 1998-06. R APPLICATION NUMBER: R FILING DATE: 1998-06. R R APPLICATION NUMBER: R FILING DATE: 1998-06. R FILING DATE: 1998-06. PR FILING DATE: 1998-05-22
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R APPLICATION NUMBER: 60/06
R FILING DATE: 1998-05-28
R APPLICATION NUMBER: 60/06
R APPLICATION NUMBER: 60/06
R FILING DATE: 1998-06-04
R FILING DATE: 1998-06-04
R APPLICATION NUMBER: 60/06 R FILING DATE: 1998-04-29
R APPLICATION NUMBER: 60/08
R FILING DATE: 1998-05-07
R APPLICATION NUMBER: 60/08
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R APPLICATION NUMBER: 60/08
R FILING DATE: 1998-05-07
R FILING DATE: 1998-05-07 R FILING DATE: 1998-04-14

RR APPLICATION NUMBER: 60/08

R FILING DATE: 1998-04-15

RR APPLICATION NUMBER: 60/08

RR FILING DATE: 1998-04-15

RR APPLICATION NUMBER: 60/08

RR FILING DATE: 1998-04-24

RR FILING DATE: 1998-04-28

RR FILING DATE: 1998-04-28

RR FILING DATE: 1998-04-28

RR APPLICATION NUMBER: 60/08

RR APPLICATION NUMBER: 60/08 R FILING DATE: 1998-03-12
R APPLICATION UMBER: 60/07
R FILING DATE: 1998-03-20
R APPLICATION NUMBER: 60/07
R FILING DATE: 1998-03-25
R APPLICATION UMBER: 60/07
R FILING DATE: 1998-02-27 REPLICATION N REPLICATION N FILING DATE: APPLICATION N FILING DATE: APPLICATION N FILING DATE: APPLICATION NUMBER: 60/0 - APPLICATION NUMBER: 60/0 - 13 - 198-05-13 APPLICATION NUMBER: FILING DATE: 1998-05 APPLICATION NUMBER: FILING DATE: APPLICATION N APPLICATION NUMBER: 60/085579 FILING DATE: 1998-05-15 FILING DATE: APPLICATION NUMBER: 60/085697 FILING DATE: 1998-01 APPLICATION NUMBER: APPLICATION NUMBER: 60/085323 FILING DATE: 1998-05-13 FILING DATE: APPLICATION NUMBER: NUMBER: 60/0: 1998-06-10 NUMBER: 60/0: 1998-06-19 NUMBER: NUMBER: NUMBER: NUMBER: NUMBER: 60/088741 1998-06-10 NUMBER: 60/085149 1998-05-12 1998-06-17 1998-06-17 1998-06-23 19/98-06-11 1998-06-10 1998-05-15 1998-05-15 .998-06-18 60/089907 60/089599 60/089532 60/088858 60/088810 60/088730 60/088026 60/087106 60/086430 60/086414 60/085338 60/084637 60/084627 60/083322 60/082999 60/085704 60/085339 60/084600 60/083545 60/081817 60/081818 60/078910 60/079663 60/079294

APPLICATION NUMBER: 60/090429

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: 60/091519
FILING DATE: 1998-07-02
APPLICATION NUMBER: 60/091982
FILING DATE: 1998-07-07
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APPLICATION NUMBER: 60/090863
FILING DATE: 1998-06-26
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FILING DATE: 1998-07-01
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APPLICATION NUMBER: 60/090445
FILING DATE: 1998-06-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CDSTGSRWRYAVPHTPGLCTSLPDPVKGTECSFSCNAGEFLDMKDQSCKPCAEGRYSLGT 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MAEPGHSHHLSARVRRRTERRIPRLWRLLLWAGTAFQVTQGTGPELHACKESEYHYEYTA 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CDSTGSRWRVAVPHTPGLCTSLSDPVKGTECSFSCNAGEFLDMKDQSCKPCAEGRYSLGT
                                                                                                                                                                                                                                     IEENTTTSFTWAFQRTTFHEASRKYTNDVAKIYSINVTNVMNGVASYCRPCALEASDVGS 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNKGETSCHQCDPDKYSEKGSSSCNVRPACTDKDYFYTHTACDANGETQLMYKWAKPKIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GIRFDEWDELPHGFASLSANMELDDSAAESTGNCTSSKWVPRGDYIASNTDECTATLMYA
                                                                                                                                                                                                                                                                                                                   PPOSVMADTENKEVARITFVFETLCSVNCELYFMVGVNSRTNTPVETWKGSKGKQSYTYI 540
                                                                                                                                                                                                                                                                                                                                                                                                                                              SEDLEGAVKLPASGVKTHCPPCNPGFFKTNNSTCQPCPYGSYSNGSDCTRCPAGTEPAVG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNKGETSCHQCDPDKYSEKGSSSCNVRPACTDKDYFYTHTACDANGETQLMYKWAKPKIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           VLYWRTTAFSVWTKVPKPVLVRNIAITGVAYTSECFPCKPGTYADKQGSSFCKLCPANSY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         VLYWRTTAFSVWTKVPKPVLVRNIAITGVAYTSECFPCKPGTYADKQGSSFCKLCPANSY
             PAELFHLESLGIPDVIFFYRSNDVTQSCSSGRSTTIRVRCSPQKTVPGSLLLPGTCSDGT 840
                                                                                                                          SRNTPTRTFNYNFSALANTVTLAGGPSFTSKGLKYFHHFTLSLCGNQGRKMSVCTDNVTD
                                                                                                                                                                                                                                                                                                  PPQSVMADTENKEVARITFVFETLCSVNCELYFMVGVNSRTNTPVETWKGSKGKQSYTYI
                                                                                                                                                                                                                                                                                                                                                               FEYKWWNTLPTNMETTVLSGINFEYKGMTGWEVAGDHIYTAAGASDNDFMILTLVVPGFR
                                                                                                                                                                                                                                                                                                                                                                                  FEYKWWNTLPTNMETTVLSGINFEYKGMTGWEVAGDHIYTAAGASDNDFMILTLVVPGFR 480
                                                                                                                                                                                                                                                                                                                                                                                                                               SEDLEGAVKLPASGYKTHCPPCNPGFFKTNNSTCQPCPYGSYSNGSDCTRCPAGTEPAVG
                                              LRIPEGESGFSKSITAYVCQAVIIPPEVTGYKAGVSSQPVSLADRLIGVTTDMTLDGITS
                                                               LRIPEGESGFSKSITAYYCQAVIIPPEVTGYKAGVSSQPVSLADRLIGVTTDMTLDGITS 780
                                                                                                           SRNTPTRTFNYNFSALANTVTLAGGPSFTSKGLKYFHHFTLSLCGNQGRKMSVCTDNVTD
                                                                                                                                                                         SCTSCPAGYYIDRDSGTCHSCPPNTILKAHQPYGVQACVPCGPGTKNNKIHSLCYNDCTF
                                                                                                                                                                                                        SCTSCPAGYYIDRDSGTCHSCPPNTILKAHQPYGVQACVPCGPGTKNNKIHSLCYNDCTF 660
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              974;
97.0%;
Similarity 99.0%;
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Pred. No. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT FILING DATE: 2002-04-12
Prior Application removed -
NUMBER OF SEQ ID NOS: 550
SEQ ID NO 38
                                                                                                                                                                                                                                                        Query Match
Best Local
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APPLICANT:
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TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C17
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ORGANISM: Homo Sapien
FEATURE:
                                                                                                                                                                                                                                                                                                                NAME/KEY: unsure
LOCATION: 877, 882
OTHER INFORMATION: unknown amino acid
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                                                           GIRFDEWDELPHGFASLSANMELDDSAAESTGNCTSSKWVPRGDYIASNTDECTATLMYA
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No. US20030022239A1
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Wood, William
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Sherwood, Steven
Smith, Victoria
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Pred. No. 0;
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                                                      APPLICANT: Baker, Kevin P.
APPLICANT: Beresini, Mauree
APPLICANT: DeForge, Laura
APPLICANT: Desnoyers, Luc
APPLICANT: Filvaroff, Ellen
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerritsen, Mary
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       APPLICANT:
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                      Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
                                              Gao, Wei-Qiang
Gerritsen, Mary E.
Goddard, Audrey
Tumas,Daniel
                                                                       Desnoyers, Luc
Filvaroff, Ellen
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      Stewart, Timothy A
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; LICATION: 877, 882
; OTHER INFORMATION: unknown amino acid
US-10-123-904-38
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Best Local Similarity 99.0
Matches 974; Conservative
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LENGTH: 1013
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APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES ITITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C54
CURRENT APPLICATION NUMBER: US/10/123,904
CURRENT FILING DATE: 2002-04-16
PILOT APPLICATION TEMOVED - See File Wrapper or Palm
NUMBER OF SEQ ID NOS: 550
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LRIPEGESGFSKSITAYVCQAVIIPPEVTGYKAGVSSQPVSLADRLIGVTTDMTLDGITS
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                                                                                                                                                                                                   ; OTHER INFORMATION: unknown US-10-140-470-38
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                                                                                                                                                                                                                                                                                          SEQ ID NO 38
LENGTH: 1013
                                                                                                                                               Matches
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C160
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LOCATION: 877, 8
                                                                                                                                                                                                                                                       FEATURE:
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            GIRFDEWDELPHGFASLSANMELDDSAAESTGNCTSSKWVPRGDYIAFNTDECTATLMYA
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                                        CDSTGSRWRVAVPHTPGLCTSLSDPVKGTECSFSCNAGEFLDMKDQSCKPCAEGRYSLGT
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
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Wood, William
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Gerritsen, Mary E.
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    APPLICANT:
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                            Gao, Wei-Qiang
Gerritsen, Mary E.
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    Godowski, Paul
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; LOCATION: 877, 882
; OTHER INFORMATION: unknown amino acid
US-10-175-746-38
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CURRENT APPLICATION NUMBER: US/10/175,746

CURRENT FILING DATE: 2002-06-19
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SCTSCPAGYYIDRDSGTCHSCPPNTILKAHQPYGVQACVPCGPGTKNNKIHSLCYNDCTF
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Tumas, Daniel
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NUMBER OF SEQ ID NOS: 550
SEQ ID NO 38
LENGTH: 1013
                                                                         Query Match
Best Local
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APPLICANT:
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                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C382
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CURRENT FILING DATE: 2002-06-20
                                                                                                                                                                                                                LENGTH: 10
TYPE: PRT
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                                                                                                                                 ORGANISM: Homo Sapien
FEATURE:
NAME/KEY: unsure
LOCATION: 877, 882
OTHER INFORMATION: unknown amino acid
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                                                                         Local Similarity
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              1 MAEPGHSHHLSARVRGRTERRIPRLWRLLLWAGTAFQVTQGTGPELHACKESEYHYEYTA
MAEPGHSHHLSARVRRRTERRIPRLWRLLLWAGTAFQVTQGTGPELHACKESEYHYEYTA
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Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A
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Wood, William
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Gerritsen, Mary E.
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                                                           Conservative
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    Mismatches

                                                                       Score 5341;
Pred. No. 0;
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                                      Sequence 38, Application US/10176921 publication No. US20030027276A1 GENERAL INFORMATION:
            APPLICANT:
                              APPLICANT: Baker, Kevin P.
APPLICANT:
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             DeForge, Laura
                      Beresini, Maureen
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: p3330R1C288
CURRENT APPLICATION NUMBER: US/10/176,921
CURRENT FILING DATE: 2002-06-20
CURRENT FILING DATE: 2002-06-20
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ORGANISM: Homo Sapien
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                      541 IEENTTTSFTWAFORTTFHEASRKYTNDVAKIYSINVTNVNNGVASYCRPCALEASDVGS
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                                                                                                                                                                                SEDLEGAVKLPASGVKTHCPPCNPGFFKTNNSTCQPCPYGSYSNGSDCTRCPAGTEPAVG
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                                                                                                                  FEYKWWNTLPTNMETTVLSGINFEYKGMTGWEVAGDHIYTAAGASDNDFMILTLVVPGFR 480
                                                                                                                                                        PPQSVMADTENKEVARITEVEETICSVNCELYFMVGVNSRTNTPVETWKGSKGKOSYTYI 540
                                                                                                    FEYKWWNTLPTNMETTYLSGINFEYKGMTGWEVAGDHIYTAAGASDNDFMILTLVVPGFR
PPQSVMADTENKEVARITEVFETICSVNCELYFMVGVNSRTNTPVETWKGSKGKQSYTYI 540
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; NAME/KEY: unsure
; LOCATION: 877, 882
; OTHER INFORMATION: unknown amino acid
US-10-137-865-38
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US-10-137-865-38
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                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/137,865
CURRENT FILING DATE: 2002-05-03
                                                                                                                                                                                                             TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC TITLE OF INVENTION: ACIDS ENCODING THE SAME FILE REFERENCE: P3330R1C154
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APPLICANT:
APPLICANT:
                                                                                                               LENGTH: 1013
TYPE: PRT
ORGANISM: Homo Sapien
                                                                                                                                                                 Prior Application removed - See Palm or File Wrapper NUMBER OF SEQ ID NOS: 550
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No. US20030032155A1
          Conservative
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                   97.0%;
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                  Score 5341;
Pred. No. 0;
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US-10-140-474-38

Sequence 38,

Publication No. US20030032156A1

Application US/10140474

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; OTHER INFORMATION: unknown amino acid US-10-140-474-38
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TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: 93330R1C162
CURRENT APPLICATION NUMBER: US/10/140,474
CURRENT FILING DATE: 2002-05-06
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LOCATION: 877, 882
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                                       SEDLEGAVKLPASGVKTHCPPCNPGFFKTNNSTCQPCPYGSYSNGSDCTRCPAGTEPAVG
             PPQSVMADTENKEVARITFVFETLCSVNCELYFMVGVNSRTNTPVETWKGSKGKQSYTYI
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NAME/RPY: unsure
LOCATION: 877, 882
OTHER INFORMATION: unknown amino acid
US-10-142-431-38
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                                                                                                        SEQ ID NO 38
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APPLICANT:
APPLICANT:
                                                                                                                                          APPLICANT: Zhang, Zemin
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: 93330R1C251
CURRENT APPLICATION NUMBER: US/10/142,431
CURRENT FILING DATE: 2002-05-10
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                                                                                                                 Prior Application removed NUMBER OF SEQ ID NOS: 550
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                                                               LENGTH: 1013
TYPE: PRT
ORGANISM: Homo Sapien
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Filvaroff, Ellen
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Godowski, Paul J.
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                                                                        CDGCNFHFLWESAAACPLCSVADYHAIVSSCVAGIQKTTYVWREPKLCSGGISLPEQRVT
                                                                                                           PAELFHLESLGIPDVIFFYRSNDVTQSCSSGRSTTIRVRCSPQKTVPGSLLLPGTCSDGT
                                                                                                                                   ICKTIDFWLKVGISAGTCTAILLTVLTCYFWKKNQKLEYKYSKLVMNATLKDCDLPAADS
                                       ICKTIDFWLKVGISAGTCTAILLTVLTCYFWKKNQKLEYKYSKLVMNATLKDCDLPAADS 960
                                                               CDGCNFHFLWESAAACPLCSVADYHAIVSSCVAGIQXTTYVXREPKLCSGGISLPEQRVT
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Pred. No. 0;
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; LOCATION: 877, 882
; OTHER INFORMATION: unknown amino acid
US-10-143-114-38
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US-10-143-114-38
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Best Local :
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APPLICANT: Baker, Kevin P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo :
FEATURE:
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CURRENT FILING DATE: 2002-05-09
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                                                                                  SNKGETSCHQCDPDKYSEKGSSSCNVRPACTDKDYFYTHTACDANGETQLMYKWAKPKIC
                                                                                                                                                           VLYWRTTAFSVWTKVPKPVLVRNIAITGVAYTSECFPCKPGTYADKQGSSFCKLCPANSY 300
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                        VLYWRTTAFSVWTKVPKPVLVRNIAITGVAYTSECFPCKPGTYADKQGSSFCKLCPANSY
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Goddard, Audrey
Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
Stewart, Timothy A.
Tumas, Daniel
Watanabe, Colin K
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Pred. No. 0;
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WOOD, ALLIAM

TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND TITLE OF INVENTION: ACIDS ENCODING THE SAME
FILE REFERENCE: P3330R1C59
CURRENT APPLICATION NUMBER: US/10/140,002
CURRENT FILING DATE: 2002-05-06
PRIOT APPLICATION TEMPORED OF SEQ ID NO 38
SEQ ID NO 38
LENGTH: 1013
TYPE: PPT

TYPE: PPT
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US-10-140-002-38
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APPLICANT: Baker, Kevin P
APPLICANT: Beresini, Mau
APPLICANT: DeForge, Laur
APPLICANT: Desnoyers, Lu
APPLICANT: Filvaroff, D
APPLICANT: Gao, Wei-Qian
APPLICANT: Gao, Wei-Qian
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Godowski, Paul J.
Gurney, Austin L.
Sherwood, Steven
Smith, Victoria
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LOCATION: 877, 882
OTHER INFORMATION: unknown amino acid
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                                                                                      PAELFHLESLGIPDVIFFYRSNDVTQSCSSGRSTTIRVRCSPQKTVPGSLLLPGTCSDGT
                                                                                                                                                                                                                                     SCTSCPAGYYIDRDSGTCHSCPPNTILKAHOPYGVOACVPCGPGTKNNKIHSLCYNDCTF 660
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ICKTIDFWLKVGISAGTCTAILLTVLTCYFWKKNQKLEYKYSKLVMNATLKDCDLPAADS
                                                        CDGCNFHFLWESAAACPLCSVADYHAIVSSCVAGIQKTTYVWREPKLCSGGISLPEORVT
                                                                                                                                   LRIPEGESGFSKSITAYVCQAVIIPPEVTGYKAGVSSQPVSLADRLIGVTTDMTLDGITS
                                                                                                                                             LRIPEGESGFSKSITAYVCQAVIIPPEVTGYKAGVSSQPVSLADRLIGVTTDMTLDGITS
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                                              CDGCNFHFLWESAAACPLCSVADYHAIVSSCVAGIQXTTYVXREPKLCSGGISLPEQRVT
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    Mismatches

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Pred. No. 0;
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  Sequence
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PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/140,584
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TITLE OF INVENTION: No. US20030032095A1el Nucleic Acid Sequences Encoding Human Semaj
FILE REFERENCE: 15966-554 CURA-54 CON-S14
CURRENT APPLICATION NUMBER: US/10/002,050
CURRENT FILING DATE: 2001-11-02
CURRENT FILING DATE: 2001-11-02
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ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  162 RGDYIAFNTDECTATLMYAVNLKQSGTVNFEYYYPDSSIIFEEFYQNDQCQP-NADDSRW 220
||:|| ||:|| :||:||:||:|| || || ||::| |||:||||| :|
| 61 RGNYIESNRDDCTVSLIYAVHLKKSGYVFFEYQYVDNNIFFEFFIQNDQCQEMDTTTDKW 120
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                                                                                                                            NSRTNTPVETWKGSKGKQSYTYIIEENTTTSFTWAFQR 555
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                                                                                                                                                                                                                                                                                                                                                                             HTACDANGETQLMYKWAKPKICSEDLEGAVKLPASGVKTHCPPCNPGFFKTNNSTCQPCP 398
                                                                                                    NRKSTNVVESWGGTKEKQAYTHIIFKNATFTFTWGIPR
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                                                                                                                                                                                                                                                                  PGTFSDGTKECRPCPAGTEPALGFEYKWWNVLPGNMKTSCFNVGNSKCDGMNGWEVAGDH 360
                                                                                                                                                                                  IQSGAGGSDNDYLILNLHIPGFKPPTS-MTGATGSELGRITFVFETLCSADCVLYFMVDI 419
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20, Application US/10002304
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Vernet, Corine
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56.1%;
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Pred. No. 1.5e-101;
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RESULT 15 US-10-003-152-20

GENERAL INFORMATION:

APPLICANT: Shimkets, Richard
APPLICANT: Fernandes, Elma
APPLICANT: Vernet, Corine
APPLICANT: Yang, Meijia
APPLICANT: Boldog, Ferenc

APPLICANT:

CANT: Herrmann, John OF INVENTION: No. US20020151494Alel Amino Acid Sequences for Human Semaphori Sequence 20, Application US/10003152 Patent No. US20020151494A1

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APPLICANT: Shimkets, Richard
APPLICANT: Fernandes, Elma
APPLICANT: Vernet, Corine
APPLICANT: Yang, Meijia
APPLICANT: Boldog, Ferenc
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PRIOR FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 49
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 20
LENGTH: 464
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Best Local Similarity
Matches 257; Conserv
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TITLE OF INVENTION: Polynucleotides and polypeptides encoded thereby
EILE REFERENCE: 15966-554 Cura-54 CON-58
CURRENT APPLICATION NUMBER: US/10/002,304
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: 09/604,286
PRIOR APPLICATION NUMBER: 09/604,286
PRIOR FILING DATE: 2000-06-22
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    420
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         162 RGDYIAFNTDECTATLMYAVNLKQSGTVNFEYYYPDSSIIFEFFVQNDQCQP-NADDSRW 220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    103 MKDQSCKPCAEGRYSLGTGIRFDEWDELPHGFASLSANMELDDSAAES-TGNCTSSKWVP 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 MKNQVCSKCGEGTYSLGSGIKFDEWDELPAGFSNIATFMDTVVGPSDSRPDGCNNSSWIP 60
                                                                                                                                                                                                                                                                    NRKSTNVVESWGGTKEKQAYTHIIFKNATFTFTWGIPR 457
                                         NSRTNTPVETWKGSKGKQSYTYIIEENTTTSFTWAFQR 555
                                                                                        IQSGAGGSDNDYLILNLHIPGFKPPTS-MTGATGSELGRITFVFETLCSADCVLYFMVDI
                                                                                                                                                                                      PGTFSDGTKECRPCPAGTEPALGFEYKWWNVLPGNMKTSCFNVGNSKCDGMNGWEVAGDH
                                                                                                                                                                                                            YGSYSNGS-DCTRCPAGTEPAVGFEYKWWNTLPTNMETTVLSGINFEYKGMTGWEVAGDH 457
                                                                                                                                    IYTAAGASDNDFMILTLVVPGFRPPQSVMADTENKEVARITFVFETLCSVNCELYFMVGV 517
                                                                                                                                                                                                                                                                                                                          HTACDANGETQLMYKWAKPKICSEDLEGAVKLPASGVKTHCPPCNPGFFKTNNSTCQPCP 398
                                                                                                                                                                                                                                                                                                                                                                       PGTFSNKPGSFNCQVCPRNTYSEKGAKECIRCKDDSQFSEEGSSECTERPPCTTKDYFQI 240
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Page 13

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FILE REFERENCE: 15966-554 CUra-54 CON-S12
CURRENT APPLICATION NUMBER: US/10/003,152
CURRENT FILING DATE: 2001-11-02
PRIOR APPLICATION NUMBER: 09/604,286
PRIOR FILING DATE: 2000-06-22
PRIOR APPLICATION NUMBER: 60/140,584
PRIOR FILING DATE: 1999-06-23
NUMBER OF SEQ ID NOS: 49
SOFTMARE: PATENTIN VET. 2.0
SEQ ID NO 20
LENGTH: 464
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        162 RGDYIAFNTDECTATLMYAVNLKQSGTVNFEYYYPDSSIIFEFFVQNDQCQP-NADDSRW 220
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                                 518 NSRTNTPVETWKGSKGKQSYTYIIEENTTTSFTWAFQR 555
                                                                  361 IOSGAGGSDNDYLILNLHIPGFKPPTS-MTGATGSELGRITFVFETLCSADCVLYFNVDI 419
                                                                                                  458 IYTAAGASDNDFMILTLYVPGFRPPÖSVMADTENKEVARITFYFETLCSVNCELYFMYGV 517
                                                                                                                                       301 PGTFSDGTKECRPCPAGTEPALGFEYKWWNVLPGNMKTSCFNVGNSKCDGMNGWEVAGDH 360
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Search completed: March 12, 2003, 04:55:23 Job time : 117 secs